

Applicants hereby further state that the information recorded in computer readable form is identical to the written sequence listing.

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Marked-up Version of Amendments:

were obtained. The integration of the DNA was analyzed with DNA prepared from tails of the rats immediately after weaning by the PCR method (primers used: tsA58-1A, 5'-TCCTAATGTGCAGTCAGGTG-3', SEQ ID NO: 1 (corresponds to 1365-1384 sites), tsA58-1B, 5'-ATGACGAGCTTTGGCACTTG-3', SEQ ID NO: 2 (corresponds to 1365-1384 sites)). As a result, 20 rats (6 male, 8 female, and 6 unknown